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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: Thu Oct 04 13:55:14 EDT 2007

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Application No: 10584486

Version No: 1.1

Input Set:**Output Set:****Started:** 2007-10-04 13:54:26.710**Finished:** 2007-10-04 13:54:28.381**Elapsed:** 0 hr(s) 0 min(s) 1 sec(s) 671 ms**Total Warnings:** 75**Total Errors:** 0**No. of SeqIDs Defined:** 85**Actual SeqID Count:** 85

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Input Set:

Output Set:

Started: 2007-10-04 13:54:26.710
Finished: 2007-10-04 13:54:28.381
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 671 ms
Total Warnings: 75
Total Errors: 0
No. of SeqIDs Defined: 85
Actual SeqID Count: 85

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

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Finnis, Christopher J. A.

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 35 40 45
 His Asp Leu Val Leu Ala Glu Phe Phe Ala Pro Trp Cys Gly His Cys
 50 55 60
 Lys Asn Met Ala Pro Glu Tyr Val Lys Ala Ala Glu Thr Leu Val Glu
 65 70 75 80
 Lys Asn Ile Thr Leu Ala Gln Ile Asp Cys Thr Glu Asn Gln Asp Leu
 85 90 95
 Cys Met Glu His Asn Ile Pro Gly Phe Pro Ser Leu Lys Ile Phe Lys
 100 105 110
 Asn Ser Asp Val Asn Asn Ser Ile Asp Tyr Glu Gly Pro Arg Thr Ala
 115 120 125

Glu	Ala	Ile	Val	Gln	Phe	Met	Ile	Lys	Gln	Ser	Gln	Pro	Ala	Val	Ala			130	135	140	
Val	Val	Ala	Asp	Leu	Pro	Ala	Tyr	Leu	Ala	Asn	Glu	Thr	Phe	Val	Thr			145	150	155	160
Pro	Val	Ile	Val	Gln	Ser	Gly	Lys	Ile	Asp	Ala	Asp	Phe	Asn	Ala	Thr			165	170	175	
Phe	Tyr	Ser	Met	Ala	Asn	Lys	His	Phe	Asn	Asp	Tyr	Asp	Phe	Val	Ser			180	185	190	
Ala	Glu	Asn	Ala	Asp	Asp	Asp	Phe	Lys	Leu	Ser	Ile	Tyr	Leu	Pro	Ser			195	200	205	
Ala	Met	Asp	Glu	Pro	Val	Val	Tyr	Asn	Gly	Lys	Lys	Ala	Asp	Ile	Ala			210	215	220	
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Phe	Gly	Glu	Ile	Asp	Gly	Ser	Val	Phe	Ala	Gln	Tyr	Val	Glu	Ser	Gly			245	250	255	
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Tyr	Lys	Pro	Leu	Phe	Thr	Glu	Leu	Ala	Lys	Lys	Asn	Arg	Gly	Leu	Met			275	280	285	
Asn	Phe	Val	Ser	Ile	Asp	Ala	Arg	Lys	Phe	Gly	Arg	His	Ala	Gly	Asn			290	295	300	
Leu	Asn	Met	Lys	Glu	Gln	Phe	Pro	Leu	Phe	Ala	Ile	His	Asp	Met	Thr			305	310	315	320
Glu	Asp	Leu	Lys	Tyr	Gly	Leu	Pro	Gln	Leu	Ser	Glu	Glu	Ala	Phe	Asp			325	330	335	
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Val	Lys	Asp	Phe	Leu	Lys	Gly	Asp	Ala	Ser	Pro	Ile	Val	Lys	Ser	Gln			355	360	365	
Glu	Ile	Phe	Glu	Asn	Gln	Asp	Ser	Ser	Val	Phe	Gln	Leu	Val	Gly	Lys			370	375	380	
Asn	His	Asp	Glu	Ile	Val	Asn	Asp	Pro	Lys	Lys	Asp	Val	Leu	Val	Leu			385	390	395	400
Tyr	Tyr	Ala	Pro	Trp	Cys	Gly	His	Cys	Lys	Arg	Leu	Ala	Pro	Thr	Tyr			405	410	415	
Gln	Glu	Leu	Ala	Asp	Thr	Tyr	Ala	Asn	Ala	Thr	Ser	Asp	Val	Leu	Ile			420	425	430	

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Gly Tyr Pro Thr Ile Val Leu Tyr Pro Gly Gly Lys Lys Ser Glu Ser
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Val Val Tyr Gln Gly Ser Arg Ser Leu Asp Ser Leu Phe Asp Phe Ile
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Lys Glu Asn Gly His Phe Asp Val Asp Gly Lys Ala Leu Tyr Glu Glu
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20 25 30

Ala Val Val Lys Leu Ala Thr Asp Ser Phe Asn Glu Tyr Ile Gln Ser
35 40 45

His Asp Leu Val Leu Ala Glu Phe Phe Ala Pro Trp Cys Gly His Cys
50 55 60

Lys Asn Met Ala Pro Glu Tyr Val Lys Ala Ala Glu Thr Leu Val Glu
65 70 75 80

Lys Asn Ile Thr Leu Ala Gln Ile Asp Cys Thr Glu Asn Gln Asp Leu
85 90 95

Cys Met Glu His Asn Ile Pro Gly Phe Pro Ser Leu Lys Ile Phe Lys
100 105 110

Asn Arg Asp Val Asn Asn Ser Ile Asp Tyr Glu Gly Pro Arg Thr Ala
115 120 125

Glu Ala Ile Val Gln Phe Met Ile Lys Gln Ser Gln Pro Ala Val Ala
130 135 140

Val Val Ala Asp Leu Pro Ala Tyr Leu Ala Asn Glu Thr Phe Val Thr
145 150 155 160

Pro Val Ile Val Gln Ser Gly Lys Ile Asp Ala Asp Phe Asn Ala Thr
165 170 175

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Ala	Glu	Asn	Ala	Asp	Asp	Asp	Phe	Lys	Leu	Ser	Ile	Tyr	Leu	Pro	Ser	195	200	205
Ala	Met	Asp	Glu	Pro	Val	Val	Tyr	Asn	Gly	Lys	Lys	Ala	Asp	Ile	Ala	210	215	220
Asp	Ala	Asp	Val	Phe	Glu	Lys	Trp	Leu	Gln	Val	Glu	Ala	Leu	Pro	Tyr	225	230	235
Phe	Gly	Glu	Ile	Asp	Gly	Ser	Val	Phe	Ala	Gln	Tyr	Val	Glu	Ser	Gly	245	250	255
Leu	Pro	Leu	Gly	Tyr	Leu	Phe	Tyr	Asn	Asp	Glu	Glu	Glu	Leu	Glu	Glu	260	265	270
Tyr	Lys	Pro	Leu	Phe	Thr	Glu	Leu	Ala	Lys	Lys	Asn	Arg	Gly	Leu	Met	275	280	285
Asn	Phe	Val	Ser	Ile	Asp	Ala	Arg	Lys	Phe	Gly	Arg	His	Ala	Gly	Asn	290	295	300
Leu	Asn	Met	Lys	Glu	Gln	Phe	Pro	Leu	Phe	Ala	Ile	His	Asp	Met	Thr	305	310	315
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Asn	His	Asp	Glu	Ile	Val	Asn	Asp	Pro	Lys	Lys	Asp	Val	Leu	Val	Leu	385	390	395
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Gln	Glu	Leu	Ala	Asp	Thr	Tyr	Ala	Asn	Ala	Thr	Ser	Asp	Val	Leu	Ile	420	425	430
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Val	Val	Tyr	Gln	Gly	Ser	Arg	Ser	Leu	Asp	Ser	Leu	Phe	Asp	Phe	Ile	465	470	475
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Val Gln Ala Asp Met Lys His Phe Pro Phe Lys Leu Ile Asp Val Asp
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Gly Lys Pro Gln Ile Gln Val Glu Phe Lys Gly Glu Thr Lys Asn Phe
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Thr Pro Glu Gln Ile Ser Ser Met Val Leu Gly Lys Met Lys Glu Thr
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Pro Ala Tyr Phe Asn Asp Ser Gln Arg Gln Ala Thr Lys Asp Ala Gly
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Thr Ile Ala Gly Leu Asn Val Leu Arg Ile Ile Asn Glu Pro Thr Ala
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 180 185 190

Leu Ile Phe Asp Leu Gly Gly Gly Thr Phe Asp Val Ser Leu Leu Phe
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